RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/763,539
Source:	TEWO
Date Processed by STIC:	11-30-04

ENTERED



IFWO

RAW SEQUENCE LISTING

DATE: 11/30/2004

PATENT APPLICATION: US/10/763,539

TIME: 12:09:55

Input Set : A:\002010-854.ST25.txt

Output Set: N:\CRF4\11302004\J763539.raw

```
4 <110> APPLICANT: Karlik, Stephen J.
         Pleiss, Michael A.
 6
         Konradi, Andrei W.
 7
         Grant, Francine S.
 8
         Semko, Christopher M.
 9
         Dressen, Darren B.
1.0
         Messersmith, Elizabeth
11
         Freedman, Stephen
         Yednock, Ted
14 <120> TITLE OF INVENTION: Composition for and Treatment of Demyelinating Diseases
15
         and Paralysis By Administration of Remyelating Agents
17 <130> FILE REFERENCE: 002010-854
19 <140> CURRENT APPLICATION NUMBER: US 10/763,539
20 <141> CURRENT FILING DATE: 2004-01-26
22 <160> NUMBER OF SEQ ID NOS: 71
24 <170> SOFTWARE: FastSEQ for Windows Version 4.0
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 360
28 <212> TYPE: DNA
29 <213> ORGANISM: Homo sapiens
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32 <221> NAME/KEY: CDS
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40 gtc aag ttg ttc tgc aca gct tct ggc ttc aac att aaa gac acc tat
41 Val Lys Leu Phe Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Thr Tyr
42
                20
44 atg cac tgg gtg aag cag agg cct caa cag ggc ctg gag tgg att gga
                                                                      144
45 Met His Trp Val Lys Gln Arg Pro Gln Gln Gly Leu Glu Trp Ile Gly
                                40
48 agg att gat cct gcg agt ggc gat act aaa tat gac ccg aag ttc cag
                                                                      192
49 Arg Ile Asp Pro Ala Ser Gly Asp Thr Lys Tyr Asp Pro Lys Phe Gln
                            55
52 gtd aag gcc act att aca gcg gad acg tcc tcc aac aca gcc tgg ctg
                                                                      240
53 Val Lys Ala Thr Ile Thr Ala Asp Thr Ser Ser Asn Thr Ala Trp Leu
54
   65
                        70
56 cag ctc age ctg aca tct gag gac act gcc gtc tac tac tgt gca
                                                                      288
57 Gln Leu Ser Ser Leu Thr Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala
                                        90
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60 gac gga atg tgg gta tca acg gga tat gct ctg gac ttc tgg ggc caa

336

RAW SEQUENCE LISTING DATE: 11/30/2004 PATENT APPLICATION: US/10/763,539 TIME: 12:09:55

Input Set: A:\002010-854.ST25.txt
Output Set: N:\CRF4\11302004\J763539.raw

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	~~~	200	200		200	ata	taa	tas	103					110			360
			acg Thr														200
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66	. 2.1.4	) - CI	115 EQ II	NTO .	2			120									
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76	var 1	ьуѕ	ьeu	GIII	5	Ser	Gry	Ala	Gru	10	val	пур	PLO	GTÀ	15	ser	
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	vaı	пув	Leu	20	Суб	1111	Ala	ser	25	PHE	ASII	rre	пув	30	TIIT	тут	
78	Mot	II i a	Пит	-	Tara	Cln	7 ~~~	Dro	-	Cln	Clar	Lou	Clu		т1о	Clar	
	Mec	птъ	Trp	vaı	пуѕ	GIII	Arg	40	GIII	GIII	Сту	цеα	45	пр	TIE	GIĀ	
80	7 200	т1.	35	Dwo	71-	Con			The	Tira	TT	7) a.s.		Tira	Dho	Cln	
	Arg		Asp	PLO	Ala	ser		Asp	TIIL	ьуѕ	TYL	60	PLO	туѕ	Pile	GIII	
82	17-1	50	70 70 70	Thr	тЪо	Thr	55	7) an	Thr	ė o m	Cor		Thr	ת ה	Trn	Lou	
		пув	Ala	1111	116	70	Ala	АБР	1111	ser	75	ASII	1111	Ala	тър	80	
84		Lou	Cor	Cor	Lou		Cor	Clu	7 cm	Thr		17-1	Trace.	Tree	Cvc		
	GIII	ьeu	Ser	ser	85	TIIL	ser	GIU	Asp	90	Ald	vai	туг	тут	95	Ala	
86	7 an	C1	Met	Пже		Cor	Thr	C1	TT 220		Tou	7 an	Dho	Tr.r.		Cln	
	Asp	Gry	мес	100	val	ser	TIIL	СТУ	105	Ala	ьеи	Asp	Pne	110	GTY	GIII	
88	C1	The	Thr		Th.	U a l	Cor	Cor	103					110			
90	GTÅ	1111	1115	vai	1111	vaı	ser.	120									
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	,		AME/F		CDS												
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			SEQUE				(310)									•	
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			2 -	, -	•	-	•					-	_		_	Gly	
105					5					10		. 100			15	_	
		a add	g gtt	acc	_	acc	t tar	. aac	a acc			a agt	atio	ract			96
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			a Trp														
113			35		. 011	. 01.		4(		, 011			45				
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117	_	- 1 y 1 5 (		· DCI		- +11	5 - 1 1 1 1 5 5 5 5 5 5 5 5 5 5 5 5 5 5		. 01)	, , , , ,	(	60 60		,		. <u>-</u> - 1	
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																Ala	2.10
121		_	, 1 Y 1	. Ory	****	. AS _E		. 4111	110	- 1111	75				. 511	80	
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RAW SEQUENCE LISTING DATE: 11/30/2004
PATENT APPLICATION: US/10/763,539 TIME: 12:09:55

Input Set : A:\002010-854.ST25.txt
Output Set: N:\CRF4\11302004\J763539.raw

100	<b>~</b> 33	a a a	ata	aa a	att	tat	ttc	tat	cac	cac	rat	tat	add	tet	ccg	tac	288
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125		пър	ДСС	1110	85	- 1 -		015		90	1101	-1-			95	-1-	
	acg	ttc	ασa	aaa		acc	aaq	cta	qaq								318
	Thr																
129			1	100	- 1		-		105								
	<210	)> SI	EO II	NO:	: 4												
	<211																
134	<212	2> T	PE:	PRT													
135	<213	3 > OI	RGAN	SM:	Homo	sar	piens	3									
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138	Ser	Ile	Val	Met	Thr	${\tt Gln}$	Thr	Pro	Lys	Phe	Leu	Leu	Val	Ser	Ala	Gly	
139	1				5					10					15		
140	Asp	Arg	Val	Thr	Ile	Thr	Cys	Lys	Ala	Ser	Gln	Ser	Val	Thr	Asn	Asp	
141				20					25					30			
142	Val	Ala	-	Tyr	Gln	Gln	Lys		Gly	Gln	Ser	Pro		Leu	Leu	He	
143			35		_	_	_	40	~ 7		_	_	45	51	m)	<b>a</b> 1	
	Tyr	-	Ala	Ser	Asn	Arg		Thr	GLY	Val	Pro		Arg	Pne	Thr	Gly	
145	~	50	m	<b>01</b>	m1	70	55	mla sa	nh a	mb .c	т1.	60	The sec	77.5	C1 n	71.	
		GIY	Tyr	GIA	Thr		Pne	TILL	Pne	THE	75	ser	1111	val	Gln	80	
147		7) are	Tou	777	77-7	70	Dho	Cvc	Gln	Gln.		ጥኒፖ	Ser	Ser	Pro		
148	Giu	Asp	Leu	Ala	85	тут	PHE	Суб	GIII	90	Азр	туг	Der	DET	95	ı yı	
	Thr	Dha	Gl v	G137		Thr	Lvc	Leu	Glu						) )		
151	1111	FIIC	Gry	100	Gry	1111	цую	пса	105	110							
	151 105 105 105 105 105 105 105 105 105																
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	<212						•										
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	9 <220> FEATURE:																
160	<22	1 > N	AME/I	KEY:	CDS												
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165	Met	Asp	Trp	Thr	Trp	Arg	Val	Phe	Cvs	T.011	T.em	$\Delta 1 =$	Val	Ala	Pro	Gly	
166	1								-1-		цси	HIU	V CL 2			-	
168					5					10					15		
	gcc	cac	tcc	cag	gtc	caa	ctg	cag	gag	10 tcc	ggṫ	gct	gaa	gtt	15 gtt	aaa	96
	gcc	cac His	tcc Ser	Gln	gtc	caa Gln	ctg Leu	cag	gag Glu	10 tcc	ggṫ	gct	gaa	gtt Val	15	aaa	96
170	gcc Ala	His	Ser	Gln 20	gtc Val	Gln	Leu	cag Gln	gag Glu 25	10 tcc Ser	ggt Gly	gct Ala	gaa Glu	gtt Val 30	15 gtt Val	aaa Lys	
170 172	gcc Ala ccg	His ggt	Ser tcc	Gln 20 tcc	gtc Val gtt	Gln aaa	Leu	cag Gln tcc	gag Glu 25 tgc	10 tcc Ser aaa	ggt Gly gct	gct Ala tcc	gaa Glu ggt	gtt Val 30 ttc	15 gtt Val aac	aaa Lys atc	96 144
170 172 173	gcc Ala ccg	His ggt	Ser tcc Ser	Gln 20 tcc	gtc Val gtt	Gln aaa	Leu	cag Gln tcc Ser	gag Glu 25 tgc	10 tcc Ser aaa	ggt Gly gct	gct Ala tcc	gaa Glu ggt Gly	gtt Val 30 ttc	15 gtt Val	aaa Lys atc	
170 172 173 174	gcc Ala ccg Pro	His ggt Gly	Ser tcc Ser 35	Gln 20 tcc Ser	gtc Val gtt Val	Gln aaa Lys	Leu ctg Leu	cag Gln tcc Ser 40	gag Glu 25 tgc Cys	10 tcc Ser aaa Lys	ggt Gly gct Ala	gct Ala tcc Ser	gaa Glu ggt Gly 45	gtt Val 30 ttc Phe	15 gtt Val aac Asn	aaa Lys atc Ile	144
170 172 173 174 176	gcc Ala ccg Pro	His ggt Gly gac	tcc Ser 35 acc	Gln 20 tcc Ser tac	gtc Val gtt Val atg	Gln aaa Lys cac	Leu ctg Leu tgg	cag Gln tcc Ser 40 gtt	gag Glu 25 tgc Cys	10 tcc Ser aaa Lys cag	ggt Gly gct Ala cgt	gct Ala tcc Ser	gaa Glu ggt Gly 45 ggt	gtt Val 30 ttc Phe	15 gtt Val aac Asn	aaa Lys atc Ile	
170 172 173 174 176 177	gcc Ala ccg Pro	ggt Gly gac Asp	tcc Ser 35 acc	Gln 20 tcc Ser tac	gtc Val gtt Val atg	Gln aaa Lys cac	Leu ctg Leu tgg Trp	cag Gln tcc Ser 40 gtt	gag Glu 25 tgc Cys	10 tcc Ser aaa Lys cag	ggt Gly gct Ala cgt	gct Ala tcc Ser ccg Pro	gaa Glu ggt Gly 45 ggt	gtt Val 30 ttc Phe	15 gtt Val aac Asn	aaa Lys atc Ile	144
170 172 173 174 176 177	gcc Ala ccg Pro aaa Lys	ggt Gly gac Asp 50	tcc Ser 35 acc Thr	Gln 20 tcc Ser tac Tyr	gtc Val gtt Val atg Met	Gln aaa Lys cac His	ctg Leu tgg Trp 55	cag Gln tcc Ser 40 gtt Val	gag Glu 25 tgc Cys aaa Lys	10 tcc Ser aaa Lys cag Gln	ggt Gly gct Ala cgt Arg	gct Ala tcc Ser ccg Pro 60	gaa Glu ggt Gly 45 ggt Gly	gtt Val 30 ttc Phe cag Gln	15 gtt Val aac Asn ggt Gly	aaa Lys atc Ile ctg Leu	144 192
170 172 173 174 176 177 178 180	gcc Ala ccg Pro aaa Lys	ggt Gly gac Asp 50 tgg	tcc ser 35 acc Thr	Gln 20 tcc Ser tac Tyr	gtc Val gtt Val atg Met	Gln aaa Lys cac His	ctg Leu tgg Trp 55 gac	cag Gln tcc Ser 40 gtt Val	gag Glu 25 tgc Cys aaa Lys	10 tcc ser aaa Lys cag Gln	ggt Gly gct Ala cgt Arg	gct Ala tcc Ser ccg Pro 60 gac	gaa Glu ggt Gly 45 ggt Gly	gtt Val 30 ttc Phe cag Gln aaa	15 gtt Val aac Asn ggt Gly tac	aaa Lys atc Ile ctg Leu	144
170 172 173 174 176 177 178 180	gcc Ala ccg Pro aaa Lys gaa Glu	ggt Gly gac Asp 50 tgg	tcc ser 35 acc Thr	Gln 20 tcc Ser tac Tyr	gtc Val gtt Val atg Met	Gln aaa Lys cac His	ctg Leu tgg Trp 55 gac	cag Gln tcc Ser 40 gtt Val	gag Glu 25 tgc Cys aaa Lys	10 tcc ser aaa Lys cag Gln	ggt Gly gct Ala cgt Arg	gct Ala tcc Ser ccg Pro 60 gac	gaa Glu ggt Gly 45 ggt Gly	gtt Val 30 ttc Phe cag Gln aaa	15 gtt Val aac Asn ggt Gly	aaa Lys atc Ile ctg Leu	144 192
170 172 173 174 176 177 178 180 181 182	gcc Ala ccg Pro aaa Lys gaa Glu 65	ggt Gly gac Asp 50 tgg Trp	tcc Ser 35 acc Thr atc	Gln 20 tcc Ser tac Tyr ggt Gly	gtc Val gtt Val atg Met cgt Arg	aaa Lys cac His atc Ile 70	ctg Leu tgg Trp 55 gac Asp	cag Gln tcc Ser 40 gtt Val ccg Pro	gag Glu 25 tgc Cys aaa Lys gct Ala	10 tcc ser aaa Lys cag Gln tcc ser	ggt Gly gct Ala cgt Arg ggt Gly 75	gct Ala tcc Ser ccg Pro 60 gac Asp	gaa Glu ggt Gly 45 ggt Gly acc Thr	gtt Val 30 ttc Phe cag Gln aaa Lys	15 gtt Val aac Asn ggt Gly tac	aaa Lys atc Ile ctg Leu gac Asp	144 192

RAW SEQUENCE LISTING DATE: 11/30/2004
PATENT APPLICATION: US/10/763,539 TIME: 12:09:55

Input Set: A:\002010-854.ST25.txt
Output Set: N:\CRF4\11302004\J763539.raw

185 Pro Lys Phe Gln Val Lys Ala Thr Ile Thr Ala Asp Glu Ser Thr Ser 188 acc get tac etg gaa etg tee tee etg egt tee gaa gae acc get gtt 336 189 Thr Ala Tyr Leu Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val 100 105 190 192 tac tac tgc get gac ggt atg tgg gtt tec acc ggt tac get etg gac 193 Tyr Tyr Cys Ala Asp Gly Met Trp Val Ser Thr Gly Tyr Ala Leu Asp 120 125 196 ttc tgg ggt cag ggt acc acg gtc acc gtc tcc tca ggt gag tcc 429 197 Phe Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Glu Ser 140 130 135 198 202 <210> SEQ ID NO: 6 203 <211> LENGTH: 143 204 <212> TYPE: PRT 205 <213> ORGANISM: Homo sapiens 207 <400> SEQUENCE: 6 208 Met Asp Trp Thr Trp Arg Val Phe Cys Leu Leu Ala Val Ala Pro Gly 210 Ala His Ser Gln Val Gln Leu Gln Glu Ser Gly Ala Glu Val Val Lys 20 212 Pro Gly Ser Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Phe Asn Ile 213 214 Lys Asp Thr Tyr Met His Trp Val Lys Gln Arg Pro Gly Gln Gly Leu 55 216 Glu Trp Ile Gly Arg Ile Asp Pro Ala Ser Gly Asp Thr Lys Tyr Asp 218 Pro Lys Phe Gln Val Lys Ala Thr Ile Thr Ala Asp Glu Ser Thr Ser 85 220 Thr Ala Tyr Leu Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val 100 105 222 Tyr Tyr Cys Ala Asp Gly Met Trp Val Ser Thr Gly Tyr Ala Leu Asp 115 120 224 Phe Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Gly Glu Ser 130 135 228 <210> SEQ ID NO: 7 229 <211> LENGTH: 383 230 <212> TYPE: DNA 231 <213> ORGANISM: Homo sapiens 233 <220> FEATURE: 234 <221> NAME/KEY: CDS 235 <222> LOCATION: (1)...(375) 237 <400> SEQUENCE: 7 238 atg ggt tgg tcc tgc atc atc ctg ttc ctg gtt gct acc gct acc ggt 239 Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly 5 10 242 gtt cac tcc atc gtt atg acc cag tcc ccg gac tcc ctg gct gtt tcc 96 243 Val His Ser Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser 25 20 246 ctg ggt gaa cgt gtt acc atc aac tgc aaa gct tcc cag tcc gtt acc

RAW SEQUENCE LISTING DATE: 11/30/2004
PATENT APPLICATION: US/10/763,539 TIME: 12:09:55

Input Set : A:\002010-854.ST25.txt
Output Set: N:\CRF4\11302004\J763539.raw

247 Leu Gly Glu Arg Val Thr Ile Asn Cys Lys Ala Ser Gln Ser Val Thr 250 aac gac gtt gct tgg tac cag cag aaa ccg ggt cag tcc ccg aaa ctg 251 Asn Asp Val Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Lys Leu 50 254 ctq atc tac gct tcc aac cgt tac acc ggt gtt ccg gac cgt ttc 255 Leu Ile Tyr Tyr Ala Ser Asn Arg Tyr Thr Gly Val Pro Asp Arg Phe 70 75 258 tee ggt tee ggt tae ggt ace gae tte ace tte ace ate tee tee gtt 288 259 Ser Gly Ser Gly Tyr Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Val 85 90 262 cag get gaa gac gtt get gtt tac tac tgc cag cag gac tac tcc tcc 336 263 Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln Asp Tyr Ser Ser 100 105 266 ccg tac acc ttc ggt ggt acc aaa ctg gag atc taa ggatcctc 383 267 Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile * 115 120 271 <210> SEQ ID NO: 8 272 <211> LENGTH: 124 273 <212> TYPE: PRT 274 <213> ORGANISM: Homo sapiens 276 <400> SEQUENCE: 8 277 Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr Gly 278 15 10 279 Val His Ser Ile Val Met Thr Gln Ser Pro Asp Ser Leu Ala Val Ser 281 Leu Gly Glu Arg Val Thr Ile Asn Cys Lys Ala Ser Gln Ser Val Thr 283 Asn Asp Val Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Lys Leu 285 Leu Ile Tyr Tyr Ala Ser Asn Arg Tyr Thr Gly Val Pro Asp Arg Phe 70 287 Ser Gly Ser Gly Tyr Gly Thr Asp Phe Thr Phe Thr Ile Ser Ser Val 85 90 289 Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys Gln Gln Asp Tyr Ser Ser 100 105 291 Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile 115 295 <210> SEQ ID NO: 9 296 <211> LENGTH: 429 297 <212> TYPE: DNA 298 <213 > ORGANISM: Homo sapiens 300 <220> FEATURE: 301 <221> NAME/KEY: CDS 302 <222> LOCATION: (1)...(429) 304 <400> SEQUENCE: 9 305 atg gac tgg acc tgg agg gtc ttc tgc ttg ctg gct gta gca cca ggt 306 Met Asp Trp Thr Trp Arg Val Phe Cys Leu Leu Ala Val Ala Pro Gly 307 1

RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 11/30/2004

PATENT APPLICATION: US/10/763,539

TIME: 12:09:56

Input Set : A:\002010-854.ST25.txt

Output Set: N:\CRF4\11302004\J763539.raw

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:69; Xaa Pos. 32,33

Seq#:70; Xaa Pos. 106,120

Seq#:71; Xaa Pos. 115

## VERIFICATION SUMMARY

DATE: 11/30/2004 TIME: 12:09:56

PATENT APPLICATION: US/10/763,539

Input Set : A:\002010-854.ST25.txt
Output Set: N:\CRF4\11302004\J763539.raw

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L:1607	M:341	W:	(46)	"n"	or	"Xaa"	used,	for	SEQ	ID#:69	after	pos.:32	
L:1646	M:341	W:	(46)	"n"	or	"Xaa"	used,	for	SEQ	ID#:70	after	pos.:96	
L:1648	M:341	W:	(46)	"n"	or	"Xaa"	used,	for	SEQ	ID#:70	after	pos.:112	
L:1677	M:341	W:	(46)	"n"	or	"Xaa"	used.	for	SEO	ID#:71	after	pos.:112	